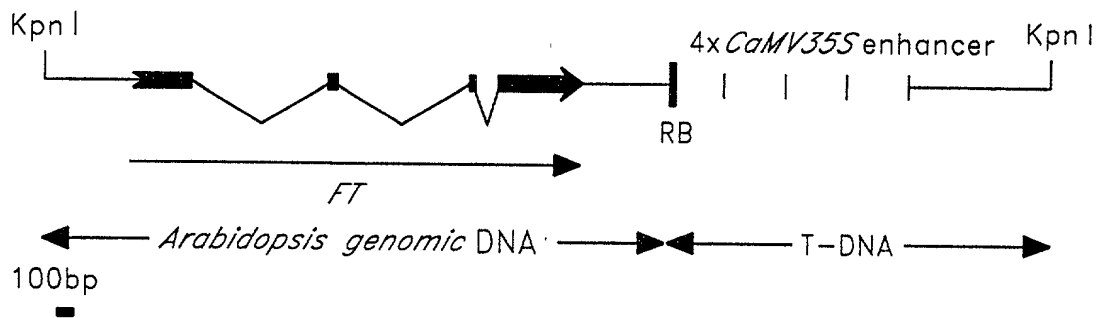
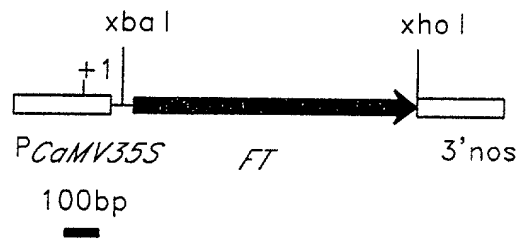


35S::FT (pSKI083)



35S::FT (pSKI059)



35S::antiFT (pSKI060)

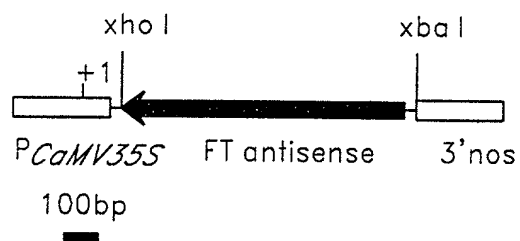


FIGURE 1

FT sense insert (814 bp without vector sequences)

Xba I

TCTAGAACTAGTGGATCCCCCGGGCTCCAGGAATTCAGCACGAGGTTTGTTC AAGATCAAAG

ATGTCTATAAATATAAGAGACCTCTTATAGTAAGCAGAGTTGTTGGAGACGTTCTTGAT
MetSerIleAsnIleArgAspProLeuIleValSerArgValValGlyAspValLeuAsp

CCGTTTAATAGATCAATCACTCTAAAGGTTACTTATGGCCAAAGAGAGGTGACTAATGGC
ProPheAsnArgSerIleThrLeuLysValThrTyrGlyGlnArgGluValThrAsnGly

TTGGATCTAAGGCCTTCTCAGGTTCAAACAAGCCAAGAGTTGAGATTGGTGGAGAAGAC
LeuAspLeuArgProSerGlnValGlnAsnLysProArgValGluIleGlyGlyGluAsp

CTCAGGAACTTCTATACTTTGGTTATGGTGGATCCAGATGTTCCAAGTCCTAGCAACCTT
LeuArgAsnPheTyrThrLeuValMetValAspProAspValProSerProSerAsnPro

CACCTCCGAGAATATCTCCATTGGTTGGTGAATGATATCCCTGCTACAACTGGAACAACC
HisLeuArgGluTyrLeuHisTrpLeuValThrAspIleProAlaThrThrGlyThrThr

TTTGGCAATGAGATTGTGTGTTACGAAAATCCAAGTCCCACTGCAGGAATTCATCGTGTC
PheGlyAsnGluIleValCysTyrGluAsnProSerProThrAlaGlyIleHisArgVal

GTGTTTATATTGTTTCGACAGCTTGGCAGGCAAACAGTGTATGCACCAGGGTGGCGCCAG
ValPheIleLeuPheArgGlnLeuGlyArgGlnThrValTyrAlaProGlyTrpArgGln

AACTTCAACACTCGCGAGTTTGCTGAGATCTACAATCTCGGCCTTCCCGTGGCCGCGAGTT
AsnPheAsnThrArgGluPheAlaGluIleTyrAsnLeuGlyLeuProValAlaAlaVal

TTCTACAATTGTCAGAGGGAGAGTGGCTGCGGAGGAAGAAGACTTTAGATGGCTTCTTCC
PheTyrAsnCysGlnArgGluSerGlyCysGlyGlyArgArgLeu***

TTTATAACCAATTGATATTGCATACTCTGATGAGATTTATGCATCTATAGTATTTTAATT
TAATAACCATTTTATGATACGAGTAACGAACGGTGATGATGCCTATAGTAGTTCAATATA
TAAGTGTGTAATAAAAATGAGAGGGGGAGGAAAATGAGAGTGTTTTACTTATATAGTGTG
TGATGCGATAATTATATTAATCTACATGAAATGAAGTGTATATTTATAAAAAAAAAAAAA

Xho I

AAAAAAACTCGAG

FT antisense sense insert (814 bp without vector sequences)

CTCGAGTTTTTTTTTTTTTTTTTTTTTTTATAAATATAACACTTCATTTTCATGTAGATTAATA
TAATTATCGCATCACACACTATATAAGTAAAACACTCTCATTTTCCTCCCCCTCTCATTT
TTATTACACACTTATATATTGAAGTACTATAGGCATCATCACCGTTTCGTTACTCGTATCA
TAAAATGGTTATTAAATTAATACTATAGATGCATAAATCTCATCAGAGTATGCAATAT
CAATTGGTTATAAAGGAAGAAGCCATCTAAAGTCTTCTTCCCTCCGCAGCCACTCTCCCTC
TGACAATTGTAGAAAACCTGCGGCCACGGGAAGGCCGAGATTGTAGATCTCAGCAAACCTCG
CGAGTGTGAAGTTCTGGCGCCACCCTGGTGCATACACTGTTTGCCTGCCAAGCTGTCTGA
AACATATAAACACGACACGATGAATTCCTGCAGTGGGACTTGGATTTTCGTAACACACA
ATCTCATTTGCCAAAGGTTGTTCCAGTTGTAGCAGGGATATCAGTCACCAACCAATGGAGA
TATTCTCGGAGGTGAGGGTTGCTAGGACTTGAACATCTGGATCCACCATAACCAAAGTA
TAGAAGTTCCTGAGGTCTTCTCCACCAATCTCAACTCTTGGCTTGTTTTGAACCTGAGAA
GGCCTTAGATCCAAGCCATTAGTCACCTCTCTTTGGCCATAAGTAACCTTTAGAGTGATT
GATCTATTAAACGGATCAAGAAGTCTCCAACAACCTCTGCTTACTATAAGAGGGTCTCTT
ATATTTATAGACATCTTTGATCTTGAACAAACCTCGTGCTGAATTCCTGCAGCCCGGGG

Xba I

ATCCACTAGTTCTAGA

FIGURE 2

[illegible]

S	I	N	I	-	-	R	D	P	L	At	FT	
E	N	M	G	T	R	V	I	E	P	L	At	TFL1
A	A	K	V	-	-	S	D	P	L	Am	CEN	
A	A	S	-	-	-	V	D	P	L	At	E12A11	
A	A	D	I	S	Q	W	A	G	P	L	Rn	HCNP
P	V	D	L	S	K	W	S	G	P	L	Hs	HCNP

3
4
5
6
7
8
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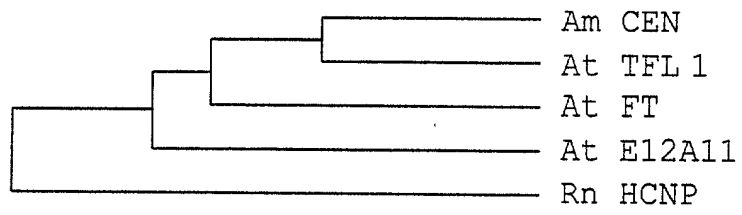


FIGURE 3B